

Claims

1. A method for distinguishing WHO classified AML subtypes AML_MLL, t(15;17), t(8;21), inv(16), 11q23, de novo_AML, s_AML, t_AML, AML_M0, AML_M1, AML_M2, AML_M4, AML_M5a, AML_M5b, AML_M6, AML_t(15;17)/M3 and/or AML_t(15;17)/M3v in a sample, the method comprising determining the expression level of markers selected from the markers identifiable by their Affymetrix Identification Numbers (affy id) as defined in Tables 1, 2, 3, 4, 5, 6 and/or 7,

wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 1.1

is indicative for the presence of AML_MLL when AML_MLL is distinguished from all other subtypes,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 1.2

is indicative for the presence of AML_inv(16) when AML_inv(16) is distinguished from all other subtypes,

and/or wherein

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 1.3 and/or

a lower expression of at least one polynucleotide defined by any of the numbers 41 of Table 1.3

is indicative for the presence of AML_other when AML_other is distinguished from all other subtypes,

and/or wherein

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a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 1.4

is indicative for the presence of AML_t(15;17) when AML_t(15;17) is distinguished from all other subtypes,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 1.5

is indicative for the presence of AML_t(8;21) when AML_t(8;21) is distinguished from all other subtypes,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 3, 4, 5, 6, 7, 11, 12, 13, 15, 16, 17, 19, 20, 22, 24, 25, 27, 28, 30, 31, 32, 33, 34, 35, 37, 41, 42, 43, 44, 46, 48, and/or 50 of Table 2.1, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 8, 9, 10, 14, 18, 21, 23, 26, 29, 36, 38, 39, 40, 45, 47, and/or 49 of Table 2.1,

is indicative for the presence of AML_MLL when AML_MLL is distinguished from AML_inv(16),

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 2.2

is indicative for the presence of AML_MLL when AML_MLL is distinguished from AML_other,

and/or wherein

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a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 5, 6, 7, 8, 9, 11, 13, 15, 18, 20, 22, 24, 25, 26, 27, 29, 30, 33, 34, 35, 36, 41, 44, 46, , and/or 50 of Table 2.3, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 4, 10, 12, 14, 16, 17, 19, 21, 23, 28, 31, 32, 37, 38, 39, 40, 42, 43, 45, 47, 48, and/or 49 of Table 2.3

is indicative for the presence of AML_MLL when AML_MLL is distinguished from AML_t(15;17),

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 8, 13, 17, 18, 19, 23, 26, 27, 28, 29, 35, 38, 39, 40, 43, 45, and/or 50 of Table 2.4, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 9, 10, 11, 12, 14, 15, 16, 20, 21, 22, 24, 25, 30, 31, 32, 33, 34, 36, 37, 41, 42, 44, 46, 47, 48, and 49 of Table 2.4,

is indicative for the presence of AML_MLL when AML_MLL is distinguished from AML_t(8;21),

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 2.5, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 37 of Table 2.5

is indicative for the presence of AML_inv(16) when AML_inv(16) is distinguished from AML_other,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 2, 4, 7, 9, 12, 17, 22, 23, 28, 29, 30, 34, 39, 42, and/or 49 of Table 2.6, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 3, 5, 6, 8, 10, 11, 13, 14, 15, 16, 18, 19, 20, 21, 24, 25, 26, 27, 31, 32, 33, 35, 36, 37, 38, 40, 41, 43, 44, 45, 46, 47, 48, and/or 50 of Table 2.6,

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is indicative for the presence of AML_inv(16) when AML_inv(16) is distinguished from AML_t(15;17),

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 6, 15, 27, 32, 36, 44, and/or 47, of Table 2.7, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 7, 8, 9, 10, 11, 12, 13, 14, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 28, 29, 30, 31, 33, 34, 35, 37, 38, 39, 40, 41, 42, 43, 45, 46, 48, 49, and/or 50 of Table 2.7,

is indicative for the presence of AML_inv(16) when AML_inv(16) is distinguished from AML_t(8;21),

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 18, and/or 25 of Table 2.8, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 19, 20, 21, 22, 23, 24, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 2.8,

is indicative for AML_other when AML_other is distinguished from AML_t(15;17),

and/or wherein

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 2.9,

is indicative for AML_other when AML_other is distinguished from AML_t(8;21),

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 4, 7, 11, 14, 16, 18, 22, 23, 24, 25, 27, 29, 31, 32, 33, 36, 37, 38, 41, 42, 43, 47, 48, and/or 49, of Table 2.10,

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a higher expression of at least one polynucleotide defined by any of the numbers 2, 3, 5, 6, 8, 9, 10, 12, 13, 15, 17, 19, 20, 21, 26, 28, 30, 34, 35, 39, 40, 44, 45, 46, and/or 50 of Table 2.10

is indicative for AML_t(15;17) when AML_t(15;17) is distinguished from AML_t(8;21),

and/or wherein

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 3.1,

is indicative for denovo_AML when denovo_AML is distinguished from all other AML subtypes

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 3.2,

is indicative for s_AML when s_AML is distinguished from all other AML subtypes,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 3.3,

is indicative for t_AML when t_AML is distinguished from all other AML subtypes

and/or wherein

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 4.1,

is indicative for denovo_AML when denovo_AML is distinguished from s_AML, and/or wherein

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a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 4.2,

is indicative for denovo_AML when denovo_AML is distinguished from t_AML and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 7, 8, 12, 13, 15, 16, 17, 19, 21, 22, 23, 24, 25, 30, 31, 34, 35, 36, 37, 38, 41, 45, 47, and/or 50 of Table 4.3, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 9, 10, 11, 14, 18, 20, 26, 27, 28, 29, 32, 33, 39, 40, 42, 43, 44, 46, 48, and/or 49 of Table 4.3,

is indicative for s_AML when s_AML is distinguished from t_AML and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 5.1,

is indicative for AML_M0 when AML_M0 is distinguished from all other AML subtypes,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 5.2,

is indicative for AML_M1 when AML_M1 is distinguished from all other AML subtypes

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 33, 34, 35, 36, 37, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 5.3, and/or

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a higher expression of at least one polynucleotide defined by any of the numbers 32 and/or 38 of Table 5.3

is indicative for AML_M2 when AML_M2 is distinguished from all other AML subtypes,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 19, 20, 21, 22, 25, 26, 27, 29, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 5.4, and/or

a higher expression a polynucleotide defined by any of the numbers 18, 23, 24, 28, and/or 30 of Table 5.4

is indicative for AML_M4 when AML_M4 is distinguished from all other AML subtypes

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 5.5,

is indicative for AML_M5a when AML_M5a is distinguished from all other AML subtypes

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 8, 9, 10, 11, 12, 13, 14, 15, 17, 18, 19, 21, 22, 23, 24, 25, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 39, 40, 42, 43, 44, 46, 48, 49, and/or 50 of Table 5.6, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 6, 7, 16, 20, 26, 37, 38, 41, 45, and/or 47 of Table 5.6,

is indicative for AML_M5b when AML_M5b is distinguished from all other AML subtypes,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28,

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29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 5.7,

is indicative for AML_M6 when AML_M6 is distinguished from all other AML subtypes,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, and/or 50 of Table 6.1, and/or

a higher expression a polynucleotide defined by any of the numbers 36, and/or 49 of Table 6.1

is indicative for AML_M0 when AML_M0 is distinguished from AML_M1,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 6.2

is indicative for AML_M0 when AML_M0 is distinguished from AML_M2,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 6.3, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 9 of Table 6.3,

is indicative for AML_M0 when AML_M0 is distinguished from AML_M4,

and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 3, 5, 7, 9, 11, 12, 14, 18, 26, 32, 33, 34, 35, 36, 39, 40, 41, 42, 43, 44, 45, 47, 48, and/or 49, of Table 6.4, and/or

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a higher expression of at least one polynucleotide defined by any of the numbers 2, 4, 6, 8, 10, 13, 15, 16, 17, 19, 20, 21, 22, 23, 24, 25, 27, 28, 29, 30, 31, 37, 38, 46, and/or 50 of Table 6.4,

is indicative for AML_M0 when AML_M0 is distinguished from AML_M5a,
and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 6.5

is indicative for AML_M0 when AML_M0 is distinguished from AML_M5b
and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 7, 8, 9, 10, 18, 26, 27, 28, 30, 32, 34, 35, 36, 37, 39, 46, 47, 48, and/or 49, of Table 6.6,
and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 11, 12, 13, 14, 15, 16, 17, 19, 20, 21, 22, 23, 24, 25, 29, 31, 33, 38, 40, 41, 42, 43, 44, 45, and/or 50 of Table 6.6

is indicative for AML_M0 when AML_M0 is distinguished from AML_M6,
and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 37, 40, 41, 42, 44, 45, 46, 47, 48, 49, and/or 50 of Table 6.7

a higher expression of at least one polynucleotide defined by any of the numbers 9, 24, 36, 38, 39, and/or 43, of Table 6.7

is indicative for AML_M1 when AML_M1 is distinguished from AML_M2,
and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 4, 5, 6, 7, 8, 9, 10, 11, 13, 15, 17, 18, 19, 21, 22, 24, 25, 26, 27, 28, 29, 30, 31, 32, 34, 35, 36, 37, 38, 40, 41, 42, 43, 44, 45, 47, 48, 49, and/or 50 of Table 6.8, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 3, 12, 14, 16, 20, 23, 33, 39, and/or 46 of Table 6.8,

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is indicative for AML_M1 when AML_M1 is distinguished from AML_M4,
and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers
23, 25, and/or 47, of Table 6.9, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1,
2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 24, 26, 27, 28, 29, 30,
31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 48, 49, and/or 50 of Table 6.9,

is indicative for AML_M1 when AML_M1 is distinguished from AML_M5a,
and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1,
2, 3, 4, 5, 7, 8, 9, 10, 11, 12, 13, 14, 15, 17, 18, 20, 22, 23, 24, 26, 28, 29, 31, 32, 33, 35,
38, 40, 41, 42, 45, 46, 48, 49, and/or 50 of Table 6.10, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 6,
16, 19, 21, 25, 27, 30, 34, 36, 37, 39, 43, 44, and/or 47 of Table 6.10

is indicative for AML_M1 when AML_M1 is distinguished from AML_M5b,
and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers
19, 22, 38, and/or 45, of Table 6.11, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1,
2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 20, 21, 23, 24, 25, 26, 27, 28, 29, 30,
31, 32, 33, 34, 35, 36, 37, 39, 40, 41, 42, 43, 44, 46, 47, 48, 49, and/or 50 of Table 6.11

is indicative for AML_M1 when AML_M1 is distinguished from AML_M6,
and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1,
2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 16, 17, 18, 20, 21, 22, 23, 24, 25, 26, 28, 29, 30, 31, 32,
33, 34, 35, 36, 37, 38, 39, 43, 45, 46, 47, 49, and/or 50 of Table 6.12, and/or

a higher expression of at least one polynucleotide defined by any of the numbers
14, 15, 19, 27, 40, 41, 42, 44, and/or 48 of table 6.12,

is indicative for AML_M2 when AML_M2 is distinguished from AML_M4,
and/or wherein

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a lower expression of at least one polynucleotide defined by any of the numbers 12 of Table 6.13, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 6.13

is indicative for AML_M2 when AML_M2 is distinguished from AML_M5a, and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 14, 15, 16, 18, 19, 20, 21, 23, 24, 25, 26, 28, 32, 33, 34, 37, 38, 39, 40, 41, 42, 43, 44, 45, 47, 49, and/or 50 of Table 6.14, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 13, 17, 22, 27, 29, 30, 31, 35, 36, 46, and/or 48 of Table 6.14,

is indicative for AML_M2 when AML_M2 is distinguished from AML_M5b, and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 26, 36, and/or 46, of Table 6.15, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 27, 28, 29, 30, 31, 32, 33, 34, 35, 37, 38, 39, 40, 41, 42, 43, 44, 45, 47, 48, 49, and/or 50 of Table 6.15

is indicative for AML_M2 when AML_M2 is distinguished from AML_M6, and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 18, 21, 25, 28, 29, 36, 40, 43, and/or 46, of Table 6.16, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 19, 20, 22, 23, 24, 26, 27, 30, 31, 32, 33, 34, 35, 37, 38, 39, 41, 42, 44, 45, 47, 48, 49, and/or 50 of Table 6.16

is indicative for AML_M4 when AML_M4 is distinguished from AML_M5a, and/or wherein

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a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 7, 9, 10, 12, 13, 15, 18, 19, 22, 25, 26, 28, 31, 32, 33, 37, 38, 40, 42, 47, and/or 50 of Table 6.17, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 5, 6, 8, 11, 14, 16, 17, 20, 21, 23, 24, 27, 29, 30, 34, 35, 36, 39, 41, 43, 44, 45, 46, 48, and/or 49 of Table 6.17,

is indicative for AML_M4 when AML_M4 is distinguished from AML_M5b, and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 39, 40, 41, and/or 47 of Table 6.18, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 42, 43, 44, 45, 46, 48, 49, and/or 50 of Table 6.18,

is indicative for AML_M4 when AML_M4 is distinguished from AML_M6, and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 2, 3, 4, 5, 6, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 23, 24, 26, 27, 28, 29, 31, 34, 35, 36, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 6.19, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 1, 7, 22, 25, 30, 32, 33, and/or 37 of Table 6.19,

is indicative for AML_M5a when AML_M5a is distinguished from AML_M5b, and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 15, 17, 18, 20, 23, 28, 29, 31, 37, 43, 44, 45, 46, and/or 48, of Table 6.20,

a higher expression of at least one polynucleotide defined by any of the numbers 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 16, 19, 21, 22, 24, 25, 26, 27, 30, 32, 33, 34, 35, 36, 38, 39, 40, 41, 42, 47, 49, and/or 50 of Table 6.20

is indicative for AML_M5a when AML_M5a is distinguished from AML_M6, and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 40, and/or 48, of Table 6.21, and/or

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a higher expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 41, 42, 43, 44, 45, 46, 47, 49, and/or 50 of Table 6.21

is indicative for AML_M5b when AML_M5b is distinguished from AML_M6, and/or wherein

a lower expression of at least one polynucleotide defined by any of the numbers 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 18, 19, 20, 21, 22, 23, 24, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, and/or 50 of Table 7, and/or

a higher expression of at least one polynucleotide defined by any of the numbers 16, 17, and/or 25, of Table 7

is indicative for AML_M3 when AML_M3 is distinguished from AML_M3v.

2. The method according to claim 1 wherein the polynucleotide is labelled.
3. The method according to claim 1, wherein the label is a luminescent, preferably a fluorescent label, an enzymatic or a radioactive label.
4. The method according to claim 1, wherein the expression level of at least two, preferably of at least ten, more preferably of at least 25, most preferably of 50 of the markers of at least one of the Tables 1-7 is determined.
5. The method according to claim 1, wherein the expression level of markers expressed lower in a first subtype than in at least one second subtype, which differs from the first subtype, is at least 5 %, 10% or 20%, more preferred at least 50% or may even be 75% or 100%, i.e. 2-fold lower, preferably at least 10-fold, more preferably at least 50-fold, and most preferably at least 100-fold lower in the first subtype.

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6. The method according to claim 1, wherein the expression level of markers expressed higher in a first subtype than in at least one second subtype, which differs from the first subtype, is at least 5 %, 10% or 20%, more preferred at least 50% or may even be 75% or 100%, i.e. 2-fold higher, preferably at least 10-fold, more preferably at least 50-fold, and most preferably at least 100-fold higher in the first subtype.

7. The method according to claim 1, wherein the sample is from an individual having AML.

8. The method according to claim 1, wherein at least one polynucleotide is in the form of a transcribed polynucleotide, or a portion thereof.

9. The method according to claim 8, wherein the transcribed polynucleotide is a mRNA or a cDNA.

10. The method according to claim 8, wherein the determining of the expression level comprises hybridizing the transcribed polynucleotide to a complementary polynucleotide, or a portion thereof, under stringent hybridization conditions.

11. The method according to claim 1, wherein at least one polynucleotide is in the form of a polypeptide, or a portion thereof.

12. The method according to claim 8, wherein the determining of the expression level comprises contacting the polynucleotide or the polypeptide with a compound specifically binding to the polynucleotide or the polypeptide.

13. The method according to claim 12, wherein the compound is an antibody, or a fragment thereof.

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14. The method according to claim 1, wherein the method is carried out on an array.

15. The method according to claim 1, wherein the method is carried out in a robotics system.

16. The method according to claim 1, wherein the method is carried out using microfluidics.

17. Use of at least one marker as defined in claim 1, for the manufacturing of a diagnostic for distinguishing WHO classified AML subtypes AML_MLL, t(15;17), t(8;21), inv(16), 11q23, de novo_AML, s_AML, t_AML, AML_M0, AML_M1, AML_M2, AML_M4, AML_M5a, AML_M5b, AML_M6, AML_t(15;17)/M3 and/or AML_t(15;17)/M3v.

18. The use according to claim 17 for distinguishing AML_MLL, t(15;17), t(8;21), inv(16), 11q23, de novo_AML, s_AML, t_AML, AML_M0, AML_M1, AML_M2, AML_M4, AML_M5a, AML_M5b, AML_M6, AML_t(15;17)/M3 and/or AML_t(15;17)/M3v in an individual having AML.

19. A diagnostic kit containing at least one marker as defined in claim 1, for distinguishing WHO classified AML subtypes AML_MLL, t(15;17), t(8;21), inv(16), 11q23, de novo_AML, s_AML, t_AML, AML_M0, AML_M1, AML_M2, AML_M4, AML_M5a, AML_M5b, AML_M6, AML_t(15;17)/M3 and/or AML_t(15;17)/M3v, in combination with suitable auxiliaries.

20. The diagnostic kit according to claim 19, wherein the kit contains a reference for the AML_MLL, t(15;17), t(8;21), inv(16), 11q23, de novo_AML, s_AML, t_AML, AML_M0, AML_M1, AML_M2, AML_M4, AML_M5a, AML_M5b, AML_M6, AML_t(15;17)/M3 and/or AML_t(15;17)/M3v subtype.

21. The diagnostic kit according to claim 20, wherein the reference is a sample or a data bank.

22. An apparatus for distinguishing AML subtype AML_MLL, t(15;17), t(8;21), inv(16), 11q23, de novo_AML, s_AML, t_AML, AML_M0, AML_M1, AML_M2, AML_M4, AML_M5a, AML_M5b, AML_M6, AML_t(15;17)/M3 and/or AML_t(15;17)/M3v in a sample containing a reference data bank.

23. The apparatus according to claim 22, wherein the reference data bank is obtainable by comprising

(a) compiling a gene expression profile of a patient sample by determining the expression level of at least one marker selected from the markers identifiable by their Affymetrix Identification Numbers (affy id) as defined in Tables 1, 2, 3, 4, 5, 6 and/or 7, and

(b) classifying the gene expression profile by means of a machine learning algorithm.

24. The apparatus according to claim 23, wherein the machine learning algorithm is selected from the group consisting of Weighted Voting, K-Nearest Neighbors, Decision Tree Induction, Support Vector Machines, and Feed-Forward Neural Networks, preferably Support Vector Machines.

25. The apparatus according to claim 22, wherein the apparatus contains a control panel and/or a monitor.

26. A reference data bank for distinguishing WHO classified AML subtype AML_MLL, t(15;17), t(8;21), inv(16), 11q23, de novo_AML, s_AML, t_AML,

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AML_M0, AML_M1, AML_M2, AML_M4, AML_M5a, AML_M5b, AML_M6,
AML_t(15;17)/M3 and/or AML_t(15;17)/M3v obtainable by comprising

- (a) compiling a gene expression profile of a patient sample by determining the expression level of at least one marker selected from the markers identifiable by their Affymetrix Identification Numbers (affy id) as defined in Tables 1, 2, 3, 4, 5, 6 and/or 7, and
- (b) classifying the gene expression profile by means of a machine learning algorithm.

27. The reference data bank according to claim 26, wherein the reference data bank is backed up and/or contained in a computational memory chip.